

AMENDMENTS TO THE SPECIFICATION

Please replace the paragraph that begins on page 10, lines 22-26, of the specification with the following amended paragraph:

Figure 13 is an alignment of the amino acid sequences of selected MATE family members FRD3 (SEQ ID NO: 3), FRDL (SEQ ID NO: 12), EDS5 (SEQ ID NO: 13), ALF5 (SEQ ID NO: 14), TT12 (SEQ ID NO: 15), ERC1 (SEQ ID NO: 16), NorM (SEQ ID NO: 17), YdhE (SEQ ID NO: 18) and DinF (SEQ ID NO: 19). Identical residues are on a black background and conservative substitutions on gray. Lines depict FRD3 transmembrane domains as predicted by HMMTOP. Multiple sequence alignments were performed using the BCM Search Launcher and residue shading using BoxShade 3.21.

Please replace the Sequence Listing with the corrected Sequence Listing submitted herewith.